

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:

10/577,293

Source:

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Date Processed by STIC:

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/577,293

DATE: 01/08/2007

TIME: 14:38:26

Input Set : A:\4276-103 Sequence Listing_ST25.txt
 Output Set: N:\CRF4\01082007\J577293.raw

3 <110> APPLICANT: GAO, Zhiqiang
 4 XIE, Hong
 5 ZHANG, Chunyan
 6 YU, Yuan Hong
 8 <120> TITLE OF INVENTION: Method for Detecting Analytes by Means of an
 Analyte/Polymeric
 9 Activator Bilayer Arrangement
 11 <130> FILE REFERENCE: 4276-103
 13 <140> CURRENT APPLICATION NUMBER: US 10/577,293
 14 <141> CURRENT FILING DATE: 2006-04-29
 16 <150> PRIOR APPLICATION NUMBER: PCT/SG04/00351
 17 <151> PRIOR FILING DATE: 2004-10-25
 19 <150> PRIOR APPLICATION NUMBER: US 60/515,229
 20 <151> PRIOR FILING DATE: 2003-10-29
 22 <160> NUMBER OF SEQ ID NOS: 14
 24 <170> SOFTWARE: PatentIn version 3.3
 26 <210> SEQ ID NO: 1
 27 <211> LENGTH: 18
 28 <212> TYPE: DNA
 29 <213> ORGANISM: Artificial
 31 <220> FEATURE:
 32 <223> OTHER INFORMATION: Primer
 34 <400> SEQUENCE: 1
 35 atggtaagg tcgggtgc
 38 <210> SEQ ID NO: 2
 39 <211> LENGTH: 20
 40 <212> TYPE: DNA
 41 <213> ORGANISM: Artificial
 43 <220> FEATURE:
 44 <223> OTHER INFORMATION: Primer
 46 <400> SEQUENCE: 2
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 50 <210> SEQ ID NO: 3
 51 <211> LENGTH: 20
 52 <212> TYPE: DNA
 53 <213> ORGANISM: Artificial
 55 <220> FEATURE:
 56 <223> OTHER INFORMATION: Primer
 58 <400> SEQUENCE: 3
 59 atggaggatt cacagtcgga
 62 <210> SEQ ID NO: 4
 63 <211> LENGTH: 18
 64 <212> TYPE: DNA
 65 <213> ORGANISM: Artificial

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67 <220> FEATURE:
 68 <223> OTHER INFORMATION: Primer
 70 <400> SEQUENCE: 4
 71 tcagtctgag tcaggccc 18
 74 <210> SEQ ID NO: 5
 75 <211> LENGTH: 35
 76 <212> TYPE: DNA
 77 <213> ORGANISM: Artificial
 79 <220> FEATURE:
 80 <223> OTHER INFORMATION: Capture probe
 82 <400> SEQUENCE: 5
 83 tttttttttt ttttactcct tggaggccat gtagg 35
 86 <210> SEQ ID NO: 6
 87 <211> LENGTH: 35
 88 <212> TYPE: DNA
 89 <213> ORGANISM: Artificial
 91 <220> FEATURE:
 92 <223> OTHER INFORMATION: Capture probe
 94 <400> SEQUENCE: 6
 95 tttttttttt ttatggtgaa ggtcggtgtc aacgg 35
 98 <210> SEQ ID NO: 7
 99 <211> LENGTH: 32
 100 <212> TYPE: DNA
 101 <213> ORGANISM: Artificial
 103 <220> FEATURE:
 104 <223> OTHER INFORMATION: Capture probe
 107 <220> FEATURE:
 108 <221> NAME/KEY: misc_feature
 109 <222> LOCATION: (1)..()
 110 <223> OTHER INFORMATION: XXX
 112 <400> SEQUENCE: 7
 113 tttttttttt ttatggagga ttcacagtcg ga 32
 116 <210> SEQ ID NO: 8
 117 <211> LENGTH: 32
 118 <212> TYPE: DNA
 119 <213> ORGANISM: Artificial
 121 <220> FEATURE:
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 124 <400> SEQUENCE: 8
 125 tttttttttt ttcagtctg agtcaggccc ca 32
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 129 <211> LENGTH: 35
 130 <212> TYPE: DNA
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 134 <223> OTHER INFORMATION: Capture probe
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 137 tttttttttt ttccctctcgc gagtcaacag aaacg 35
 140 <210> SEQ ID NO: 10

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143 <213> ORGANISM: Artificial
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146 <223> OTHER INFORMATION: Capture probe
148 <400> SEQUENCE: 10
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152 <210> SEQ ID NO: 11
153 <211> LENGTH: 32
154 <212> TYPE: DNA
155 <213> ORGANISM: Artificial
157 <220> FEATURE:
158 <223> OTHER INFORMATION: Detection probe
160 <400> SEQUENCE: 11
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164 <210> SEQ ID NO: 12
165 <211> LENGTH: 71
166 <212> TYPE: DNA
167 <213> ORGANISM: Artificial
169 <220> FEATURE:
170 <223> OTHER INFORMATION: Oligonucleotide
172 <400> SEQUENCE: 12
173 aaattgaaga gtttgatcat ggctcagatt gaacgctggc aaaaaaaaaa aaaactccta 60
175 cgggagggcag c 71
178 <210> SEQ ID NO: 13
179 <211> LENGTH: 71
180 <212> TYPE: DNA
181 <213> ORGANISM: Artificial
183 <220> FEATURE:
184 <223> OTHER INFORMATION: Oligonucleotide
186 <400> SEQUENCE: 13
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189 cgggagggcag c 71
192 <210> SEQ ID NO: 14
193 <211> LENGTH: 71
194 <212> TYPE: DNA
195 <213> ORGANISM: Artificial
197 <220> FEATURE:
198 <223> OTHER INFORMATION: Oligonucleotide
200 <400> SEQUENCE: 14
201 aaattgaaga gtatgatcat gtctcagatt gaacgctggc aaaaaaaaaa aaaactccta 60
203 cgggagggcag c 71

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Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1,2,3,4,5,6,7,8,9,10,11,12,13,14

VERIFICATION SUMMARY

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